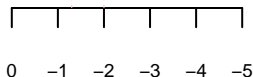
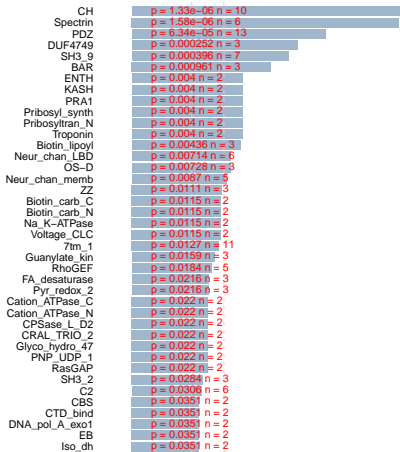


OeCa_A domain



log(p)

n = 695 genes | p adjust BH FALSE